IN THE CLAIMS

Please amend claims 1, 5, 10 and 12 as follows.

This listing of the claims will replace all prior versions, and listings, of claims in the application.

1. (Currently Amended) An isolated cDNA, or the complement thereof, comprising a nucleic acid sequence encoding a protein selected from the group consisting of:

a) an amino acid sequence of SEQ ID NO:1; and

b) a naturally occurring variant of the amino acid sequence of SEQ ID NO:1 having at least 90% identity to the amino acid sequence of SEQ ID NO:1[[;]]

~~c) a biologically active fragment of SEQ ID NO:1 from about amino acid residue L182 to about amino acid residue K214 of SEQ ID NO:1; and~~

~~d) an antigenic epitope of SEQ ID NO:1 from about amino acid residue F136 to about amino acid residue L154 of SEQ ID NO:1~~.

2. (Original) An isolated cDNA comprising a nucleic acid sequence selected from:

a) SEQ ID NO:2 or the complement thereof;

b) a fragment of SEQ ID NO:2 selected from SEQ ID NOs:3-6 or the complements thereof; and

c) a variant of SEQ ID NO:2 selected from SEQ ID NOs:7-10.

3. (Original) A composition comprising the cDNA or the complement of the cDNA of claim 1 and a labeling moiety.

4. (Original) A vector comprising the cDNA of claim 1.

5. (Currently Amended) [[A]] An isolated host cell comprising the vector of claim 4.

6. (Original) A method for using a cDNA to produce a protein, the method comprising:

a) culturing the host cell of claim 5 under conditions for protein expression; and

b) recovering the protein from the host cell culture.

7. (Original) A method for using a cDNA to detect expression of a nucleic acid in a sample comprising:

a) hybridizing the composition of claim 3 to nucleic acids of the sample, thereby forming hybridization complexes; and

b) comparing hybridization complex formation with a standard, wherein the comparison indicates expression of the cDNA in the sample.

8. (Original) The method of claim 7 further comprising amplifying the nucleic acids of the sample prior to hybridization.

9. (Original) The method of claim 7 wherein the composition is attached to a substrate.

10. (Currently Amended) The method of claim 7 wherein [[the]] increased expression of the cDNA in the sample ~~is differentially expressed~~ when compared with a standard of normal tissue is [[and]] diagnostic of clear cell sarcoma.

11. (Previously Amended) A method of using a cDNA to screen a plurality of molecules or compounds for a molecule or compound that specifically binds the cDNA, the method comprising:

a) combining the cDNA of claim 1 with a plurality of molecules or compounds under conditions to allow specific binding; and

b) detecting specific binding, thereby identifying a molecule or compound which specifically binds the cDNA.

12. (Currently Amended) The method of claim 11 wherein the molecules or compounds are selected from DNA molecules, RNA molecules, ~~peptide nucleic acids,~~ artificial chromosome constructions, peptides, transcription factors, and repressors[[,]] ~~and regulatory molecules~~.